STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

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form:

Application Serial Number: _10/541,998_

Source: _IFWP_

Date Processed by STIC: _8/3/06_

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2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A
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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT
MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER
VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND
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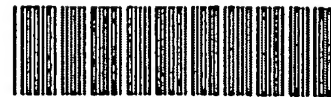   http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is
a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the
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1. EFS-Bio (<**http://www.uspto.gov/ebc/efs/downloads/documents.htm>** , **EFS Submission
   User Manual** - ePAVE)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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   Alexandria, VA 22314

Revised 01/10/06

IFWP

RAW SEQUENCE LISTING   DATE: 08/03/2006
PATENT APPLICATION: US/10/541,998   TIME: 09:50:26

Input Set : A:\19020.seq.txt
Output Set: N:\CRF4\08032006\J541998.raw

```
 3 <110> APPLICANT: Collier, Greg
 4        Walder, Ken
 5        Segal, David
 6        Foletta, Victoria
 8 <120> TITLE OF INVENTION: OBESITY-RELATED GENES
10 <130> FILE REFERENCE: 19020
12 <140> CURRENT APPLICATION NUMBER: 10/541,998
13 <141> CURRENT FILING DATE: 2005-07-13
15 <160> NUMBER OF SEQ ID NOS: 13
17 <170> SOFTWARE: PatentIn version 3.1
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 426
21 <212> TYPE: DNA
22 <213> ORGANISM: Psammomys obesus
24 <220> FEATURE:
25 <221> NAME/KEY: misc_feature
26 <222> LOCATION: (336)..(366)
27 <223> OTHER INFORMATION: n = any nucleotide
30 <400> SEQUENCE: 1
31 ggacatcttt tcagccatga ggagctttct ggaaactcgg agttgataca gaaatatagg    60
33 aatataatca cgcaggctcc taacctggag aacattgagc tgtactggaa cagctacaac   120
35 aaccgccgag acctgaactt cgacgcaggt ggtgagatga ccctcaagtg ccctgtgatg   180
37 ctggtggtag gagaccaagc gcctcatgag gatgccgtgg tggagtgtaa ctcaaaactg   240
39 gaccccacac agacctcgtt cctcaagatg gctgattctg gaggtcagcc acagctgacc   300
41 cagccaggca agctgactga ggctttcaag tacttnctgc aaggcatggg ctacatggcc   360
43 tcctcctgca tgactcgcct atcgaggtct cgcacggcat ctttgaccag cgcagcatcc   420
45 attgat                                                            426
48 <210> SEQ ID NO: 2
49 <211> LENGTH: 451
50 <212> TYPE: DNA
51 <213> ORGANISM: Psammomys obesus
53 <400> SEQUENCE: 2
54 gctggtaccg gtccggaatt cccgggatat cgtcgaccca cgcgtccggt ggtggagaag    60
56 atcgctcctg ccgtggttca cattgaactg tatcgcaaac ttcctttctc gaagagggag   120
58 gtgccagtgg cgagtgggtc cggatttatc gtgtctgagg atggactgat tgtgaccaat   180
60 gctcacgtgg tgaccaacaa aaacagggtc aaggttgagc tgaagaatgg agcaacctat   240
62 gaagctaaaa tcaaggatgt ggatgaaaag gcagacatcg cacttatcaa aattgaccac   300
64 cagggaaagc tgccagtctt gctgctgggc cgctcctcag agcttcgacc aggagagttt   360
66 gtggtcgcca tcggaagccc ctttttcctt caaaacacag tcaccactgg gatcgtcagt   420
68 accacccagc gaggcggcaa agagctgggg c                                 451
71 <210> SEQ ID NO: 3
72 <211> LENGTH: 478
73 <212> TYPE: DNA
```

W--> (annotation pointing to line 41)

Handwritten annotations:

see pp 1-7

Does Not Comply
Corrected Diskette Needed

(336) — circled (366) — (336)

OTHER INFORMATION: n = any nucleotide (sequence circled)

FYI: "n" can only represent a single nucleotide. If "sequence" denotes more than one nucleotide, it is an invalid designation. If you mean "n" = any nucleotide, that is valid.

(please check all sequences showing n's)

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Input Set : A:\19020.seq.txt
Output Set: N:\CRF4\08032006\J541998.raw

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74 <213> ORGANISM: Psammomys obesus
77 <400> SEQUENCE: 3
78 tgacattttc tttccacctc ttatgatagc tgatatatac taaatcttta tacagaaatg      60
80 tcagtacttg aacaaattca aaacacattg gtttattaac ttttggctca tgcatggttt     120
82 attaggttca aattatacct gattcatcta tatttacttt taaaatgtgt ggtttcctca     180
84 ttttaaaagt aaaactaaac agtgcttttg gaatttctaa gctactaatt gttgatagat     240
86 acagcctgtg tctagtaaaa tagttttgtg ggtgtgggtt ctatctttcc atgaaaaagt     300
88 gggaggtgta agttagtttg gttagtgcct aatagttaaa tttatataaa ataagaatga     360
90 gcatttggta tctgtatgaa agggccctaa atcaaaatga ttatccataa tcaatcttta     420
92 ttcttgtttt ataaaaacca aagggcactc attggttaag tgtgctgaga tagaaaag      478
95 <210> SEQ ID NO: 4
96 <211> LENGTH: 1884
97 <212> TYPE: DNA
98 <213> ORGANISM: Psammomys obesus
100 <220> FEATURE:
101 <221> NAME/KEY: misc_feature
102 <222> LOCATION: (609)..(609)
103 <223> OTHER INFORMATION: n = any nucleotide sequence
106 <220> FEATURE:
107 <221> NAME/KEY: misc_feature
108 <222> LOCATION: (611)..(611)
109 <223> OTHER INFORMATION: n = any nucleotide sequence
113 <220> FEATURE:
114 <221> NAME/KEY: misc_feature
115 <222> LOCATION: (1388)..(1388)
116 <223> OTHER INFORMATION: n = any nucleotide sequence
119 <220> FEATURE:
120 <221> NAME/KEY: misc_feature
121 <222> LOCATION: (1390)..(1390)
122 <223> OTHER INFORMATION: n = any nucleotide sequence
125 <220> FEATURE:
126 <221> NAME/KEY: misc_feature
127 <222> LOCATION: (1413)..(1413)
128 <223> OTHER INFORMATION: n = any nucleotide sequence
131 <220> FEATURE:
132 <221> NAME/KEY: misc_feature
133 <222> LOCATION: (1512)..(1512)
134 <223> OTHER INFORMATION: n = any nucleotide sequence
137 <220> FEATURE:
138 <221> NAME/KEY: misc_feature
139 <222> LOCATION: (1568)..(1568)
140 <223> OTHER INFORMATION: n = any nucleotide sequence
143 <400> SEQUENCE: 4
144 cgcagatact gcagagctga cgcattcttt gcctggcatc tcagcttgct aggtgtctca      60
146 tcttcgctct gcgctcgtgg cctcctccaa ggcctccagt ctcctttaag aaaactcaag     120
148 acctgggaag ctacgatgcg agcttgatgc cgctacccta gcaggctatg gacttcctga     180
150 gggtcctcgg actgttgaca cccattccga tccgcatcct tccaagctga taagcccggg     240
152 accctagggc ggggtgccca gactcatgtg tgacgccttg cagtgaaacc ccattcccag     300
154 tgtggttgct tctttgctgg gctttggccc atttgatacc acgaaggatg acgatgctag     360
```

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Input Set : A:\19020.seq.txt
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```
      156 ttatgcagca gccaacacac ctcccccccaa ctctccgctg tcactggtgg gccccactgt    420
      158 ccaggaagca ggtgtccgga actgacatct tggagcagag gggccatgag aggtgtgtgt    480
      160 atcctgccag aaagcagctg gaccacgacg ctcccaagat gaacccactg tatacagagg    540
      162 catcatggga gttgttatgt caggagcatt ctagacccac gtgtacttga gcgtggaaag    600
W--> 164 acagaagana ngcgcagaga ctggggcact tgatctgctc accatgatcg cctgcacggg    660
      166 tctcatccag ttcctgcctt aggctacagt ggcggtgtcc acgggcttgc cattcaacgt    720
      168 gctctcagac ccagatcggg ctcaccactg aggagaacct tttcactttg gtgggtatgc    780
      170 agagggaagg gtctcgactc cagagacctg gagtccaaag tttgtttttt tttatttacc    840
      172 agtaatttat ttattttttt tattattagt tatgagtcca aagtttaata ccattcaagc    900
      174 tactgctgtc tgtctgctta gagccacagc atgcaatgtg gcaccaaggc atccttgtcc    960
      176 tcacagtttc acactgtggg aacaggcatc cttgttctta cagattagcg cgagggaaac   1020
      178 cagaaatatt aaacacgcag ggttgtctct ccaaagggag aggcacatac cctgtttttcc   1080
      180 tcccgaaggc tgggagcgga ggtgtttgat atcctggcta cctctgtgca atctgtaggc   1140
      182 catgtcctta agatgtagct gtcagtcggt agtggagccg gagccgtcag tcagtagatt   1200
      184 ggggttgtgg catgcgcctt taactccatt taattccagc actctagtgg tttggtacag   1260
      186 cagcagcagc agcggttgca gtggcccggg gaagtcctga agaccagctt tcatcccagc   1320
      188 actcaggcag caaaggcggg tggattctct gagttcgagg ccggcctggt ctacagagtg   1380
      190 agtccagncn agccagggct acaacagaga aancctctct attgaaaaat aaataaatta   1440
      192 taaaaaaaaa aaggtgtcat gtgtcctgtg tactttacaa agaatgttga tgcttaagct   1500
      194 tttttgtgca cncaagaaaa ttgtttaact ggtgtcagac tcctgaagtt tgaaccagca   1560
      196 cttagccngg cgtggtggcg cacgcctgta atcccagccc tcgggaggca gaggcaggtg   1620
      198 gatctctgag ttagaggcca gcctggtcta cagagtgagt ccaggacagc caggattaca   1680
      200 cagagaaacc ccgtctcaaa aatgtaaaat aaattaaaat aaagtttgaa ccaacagtgt   1740
      202 ttactgagtc gtgttgaaac agattacctt tttgcttctc tttgatcatt attctactgt   1800
      204 ggtgtcagca gagaccccctc cagcaggtgg ccaacgtgag agtctcaagc ccgagaaggt   1860
      206 aagaatttaa aaaaaaaaaa aaaa                                          1884
      209 <210> SEQ ID NO: 5
      210 <211> LENGTH: 1358
      211 <212> TYPE: DNA
      212 <213> ORGANISM: Psammomys obesus
      214 <220> FEATURE:
      215 <221> NAME/KEY: misc_feature
      216 <222> LOCATION: (385)..(385)
      217 <223> OTHER INFORMATION: n = any nucleotide sequence
      220 <220> FEATURE:
      221 <221> NAME/KEY: misc_feature
      222 <222> LOCATION: (416)..(416)
      223 <223> OTHER INFORMATION: n = any nucleotide sequence
      227 <220> FEATURE:
      228 <221> NAME/KEY: misc_feature
      229 <222> LOCATION: (430)..(430)
      230 <223> OTHER INFORMATION: n = any nucleotide sequence
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      234 <221> NAME/KEY: misc_feature
      235 <222> LOCATION: (507)..(507)
      236 <223> OTHER INFORMATION: n = any nucleotide sequence
      239 <220> FEATURE:
      240 <221> NAME/KEY: misc_feature
      241 <222> LOCATION: (754)..(754)
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RAW SEQUENCE LISTING   DATE: 08/03/2006
PATENT APPLICATION: US/10/541,998   TIME: 09:50:26

Input Set : A:\19020.seq.txt.
Output Set: N:\CRF4\08032006\J541998.raw

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242 <223> OTHER INFORMATION: n = any nucleotide sequence
245 <220> FEATURE:
246 <221> NAME/KEY: misc_feature
247 <222> LOCATION: (856)..(856)
248 <223> OTHER INFORMATION: n = any nucleotide sequence
251 <220> FEATURE:
252 <221> NAME/KEY: misc_feature
253 <222> LOCATION: (928)..(928)
254 <223> OTHER INFORMATION: n = any nucleotide sequence
257 <220> FEATURE:
258 <221> NAME/KEY: misc_feature
259 <222> LOCATION: (980)..(980)
260 <223> OTHER INFORMATION: n = any nucleotide sequence
264 <220> FEATURE:
265 <221> NAME/KEY: misc_feature
266 <222> LOCATION: (1165)..(1165)
267 <223> OTHER INFORMATION: n = any nucleotide sequence
270 <220> FEATURE:
271 <221> NAME/KEY: misc_feature
272 <222> LOCATION: (1200)..(1200)
273 <223> OTHER INFORMATION: n' = any nucleotide sequence
276 <220> FEATURE:
277 <221> NAME/KEY: misc_feature
278 <222> LOCATION: (1273)..(1273)
279 <223> OTHER INFORMATION: n = any nucleotide sequence
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283 <221> NAME/KEY: misc_feature
284 <222> LOCATION: (1284)..(1284)
285 <223> OTHER INFORMATION: n = any nucleotide sequence
288 <220> FEATURE:
289 <221> NAME/KEY: misc_feature
290 <222> LOCATION: (1303)..(1303)
291 <223> OTHER INFORMATION: n = any nucleotide sequence
294 <220> FEATURE:
295 <221> NAME/KEY: misc_feature
296 <222> LOCATION: (1305)..(1305)
297 <223> OTHER INFORMATION: n = any nucleotide sequence
301 <220> FEATURE:
302 <221> NAME/KEY: misc_feature
303 <222> LOCATION: (1326)..(1326)
304 <223> OTHER INFORMATION: n = any nucleotide sequence
307 <220> FEATURE:
308 <221> NAME/KEY: misc_feature
309 <222> LOCATION: (1330)..(1330)
310 <223> OTHER INFORMATION: n = any nucleotide sequence
313 <400> SEQUENCE: 5
314 gttgggaaag aatgaagaaa caacccgatg aatagaaatg aaaagcctaa gccaaatgga      60
316 tttctgttga gatgttggat gaaaacaagt atccactgtt taccaacttg acgaaaaatc     120
318 tcaactgagg tttggctgtt aaaaaaaaaa attcactgtg gcctctgtgc ttaattgtcg     180
```

RAW SEQUENCE LISTING   DATE: 08/03/2006
PATENT APPLICATION: US/10/541,998   TIME: 09:50:26

Input Set : A:\19020.seq.txt
Output Set: N:\CRF4\08032006\J541998.raw

```
        320 taaaccattg tgactgttac tgctcaaagt atcgtactgt tcattagtaa ctacatcaga    240
        322 attgcaccgc tgctgttgga aaagccaata aagaaacccc cagactgctg ctcagcaaat    300
        324 gttaataaag tgtgcgcacc gtaggcctgt ccacccagtc accaagcagc gtccctttgt    360
W-->    326 ctgcgagtgg ctgtgggtgt gattnaccac ctcagaggtg cacagcacct gcttgngccc    420
        328 ttaagtgtgn gtcagaagac aagcagcttc tcggtaacca acaacctgct tttcggagct    480
        330 cagtgtttag gctgtttact gaatcanata tgtaactcag cacacataag cgaagagaga    540
        332 ttttggctgc actggcaaga gtgaaccaaa tttacttcta tttttaaag gcagatcata    600
        334 tttaagcata taagtaattt atggatataa attgttggat atttatttta gtctgaatat    660
        336 ttgttttaa attattacat gtgttctcta tgtctttatc tctggaataa cgatgccatt    720
        338 aaccacatgg ccatatgttt tgaaagttgg gtgnaacaga ggaaaagtca tccttcttgg    780
        340 ttcttgactc cctttcctca actacatgat aagtctatca ataaagcatt tgacctcagc    840
        342 aggggcagaa gcctgnaaag ttagaaaact cattgaccac agtagacaat tgatttctta    900
        344 gaaataagaa gtgagaagca gctgctgngc tgagcagggg atgtaaacca agtccagatg    960
        346 caccaacgtg aagaggcttn tagcaaaaat atgtttgcct ctcacccctg cacatgttct   1020
        348 agatgcttaa aaacagccac atggccccgc tgcgaggacc tcgtaatgtt gttgttgttg   1080
        350 ttgttgttgt taaaggagtt ctcacaagcg tacaagtgca gcactgaaag tggctgaggc   1140
        352 ccacagtcct cagcacccaa gtctnttccg cagcacgcca agctggtgtt gtccgggtgn   1200
        354 gtatgtctgt gctcagtgcc aagctggtgt ttggtcccgt tgtatattat gtgccccaag   1260
        356 tgtttgggg canagctgac ccangctgga cacacttctt ttngncttcg agtttactgg   1320
        358 ttgatncagn taaaaataaa ttaattaatt aaagactt                            1358
        361 <210> SEQ ID NO: 6
        362 <211> LENGTH: 390
        363 <212> TYPE: DNA
        364 <213> ORGANISM: Psammomys obesus
        366 <220> FEATURE:
        367 <221> NAME/KEY: misc_feature
        368 <222> LOCATION: (3)..(3)
        369 <223> OTHER INFORMATION: n = any nucleotide sequence
        372 <220> FEATURE:
        373 <221> NAME/KEY: misc_feature
        374 <222> LOCATION: (9)..(9)
        375 <223> OTHER INFORMATION: n = any nucleotide sequence
        378 <220> FEATURE:
        379 <221> NAME/KEY: misc_feature
        380 <222> LOCATION: (60)..(60)
        381 <223> OTHER INFORMATION: n = any nucleotide sequence
        384 <220> FEATURE:
        385 <221> NAME/KEY: misc_feature
        386 <222> LOCATION: (71)..(71)
        387 <223> OTHER INFORMATION: n = any nucleotide sequence
        390 <220> FEATURE:
        391 <221> NAME/KEY: misc_feature
        392 <222> LOCATION: (91)..(91)
        393 <223> OTHER INFORMATION: n = any nucleotide sequence
        396 <220> FEATURE:
        397 <221> NAME/KEY: misc_feature
        398 <222> LOCATION: (113)..(113)
        399 <223> OTHER INFORMATION: n = any nucleotide sequence
        402 <220> FEATURE:
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(handwritten annotations) ? "c" is at 60 (see p.6) "a" is at 71 "a" is at 91 "g" is at 113 see p.6

<220>
<221> misc_feature   *c* is at 388
<222> (388)..(388)
<223> n = any nucleotide sequence

<220>
<221> misc_feature   there are only 390
<222> (524)..(524)   nucleotides in this
<223> n = any nucleotide sequence   sequence

<220>
<221> misc_feature
<222> (527)..(527)
<223> n = any nucleotide sequence

<400> 6

```
gtngaagcnt aggagttcga ggatgcgccc gatgtcgagc cgctggaacc cacgcttagc     60

aatatcatcg agcagcgcag ccttaagtgg accttcgtcg ggggcaaggg tggcgttggt    120

aagaccacct gcagctgcag cctggcggtc cagctgtcta agggacgtga gagtgttcta    180

atcatttcca cagacccagc tcacaacatc tcagatgcat ttgaccagaa gttctccaag    240

gtgcctacca aggtcaaagg ctatgacaac ctctttgcta tggagataga cccgagcctg    300

ggcgtggcag agctccctga tgaagttctt cgaggaagac aacatgctga gcatgggcaa    360

gaagatgatg caggaggcca tgagcgcctt                                     390
```

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 08/03/2006
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Input Set : A:\19020.seq.txt
Output Set: N:\CRF4\08032006\J541998.raw

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 336
Seq#:4; N Pos. 609,611,1388,1390,1413,1512,1568
Seq#:5; N Pos. 385,416,430,507,754,856,928,980,1165,1200,1273,1284,1303
Seq#:5; N Pos. 1305,1326,1330
Seq#:6; N Pos. 3,9
Seq#:7; N Pos. 524,527
Seq#:8; N Pos. 60
Seq#:9; N Pos. 71,91,113,388

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/541,998

DATE: 08/03/2006
TIME: 09:50:27

Input Set : A:\19020.seq.txt
Output Set: N:\CRF4\08032006\J541998.raw

```
L:41 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:300
L:164 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:600
M:341 Repeated in SeqNo=4
L:326 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:360
M:341 Repeated in SeqNo=5
L:422 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:471 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:480
L:501 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:60
M:341 Repeated in SeqNo=9
```